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1.

Mastering seeds for genomic size nucleotide BLAST searches

For example, the **seed** with **template** 110011 can start any **alignment** that contains ... We,

therefore, imple mented the **dynamic programming** algorithm for exact hit ...

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2.

Mastering seeds for genomic size nucleotide BLAST searches -- Gotea et ...

Email this **article** to a friend. Similar **articles** in this journal ... For example, the **seed** with **template**

110011 can start any **alignment** that contains ...

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3.

GMAP: a genomic mapping and alignment program for mRNA and EST ...

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the resulting exon cores' through **dynamic programming** (DP) ...

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4.

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independently **using the dynamic programming**. algorithm. 28,30 ... the peptide backbone  
corresponding to deletions in **the alignment between** GSIII ...

[sbio.uct.ac.za/Sbio/research\\_outputs/articles/VanRooyen\\_2006.pdf](http://sbio.uct.ac.za/Sbio/research_outputs/articles/VanRooyen_2006.pdf)

5.

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gene to the gene names in **the articles** most similar to a given **article** [22, 24] ... **dynamic**  
**programming** algorithm to maximize the matched number of letters **between** ...

[www.ohsu.edu/dmice/people/students/theses/2005/upload/Cohen-thesis.pdf](http://www.ohsu.edu/dmice/people/students/theses/2005/upload/Cohen-thesis.pdf)

6.

[A Rural Studies Bibliography](#)

... contrasts this to the nurturing and harmonious relation **between the** small ... be social **alignment**  
against the interest of traditional elites and their ...

[www.nal.usda.gov/ric/ricpubs/ruralbib.html](http://www.nal.usda.gov/ric/ricpubs/ruralbib.html) - 282k - [Cached](#)

7.

[PDF] [Part 11 of ARDA 2 SECTION THREE Volume Eighteen 2002 c.e. Drynemetum Press](#)

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mention fun, to actually visit all the RDNA **and related** groves I. could find contacts for. ... the  
Sundays **closest to the** new and full moons in addition to the ...

[orgs.carleton.edu/Druids/ARDA2/doc/2part11-3b1.pdf](http://orgs.carleton.edu/Druids/ARDA2/doc/2part11-3b1.pdf)

8.

[people.csail.mit.edu](http://people.csail.mit.edu)

... co uk carries archives moderated **articles** files information send saying help ... mathew **article** ba

drporter suvm syr edu brad porter writes science wonderful ...

[people.csail.mit.edu/jrennie/20Newsgroups/vocabulary.txt](http://people.csail.mit.edu/jrennie/20Newsgroups/vocabulary.txt) - 494k - [Cached](#)

9.

[Bioinformatics citation list for keyword 'genome assembly'](#)

Bioinformatics **articles** relating to 'genome assembly' ... The program then identifies conserved

sequence tags ( ... like **alignment between the** query and ...

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10.

[Bioinformatics citation list for keyword 'proteins'](#)

... residue in a **related** protein, and the local similarity **between the** two sequences. ... of

**dynamic** protein binding to chromatin in vivo, **using** photobleaching ...

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